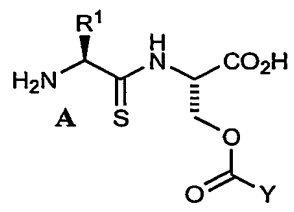
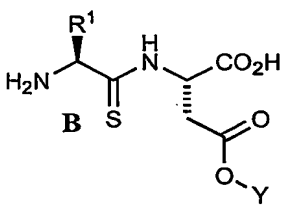
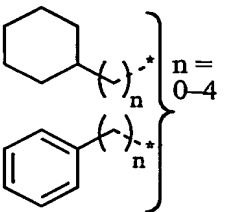
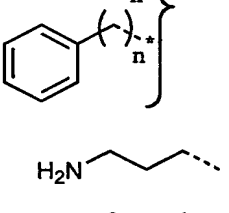
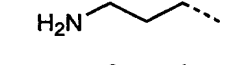
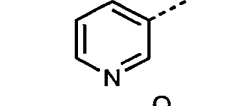
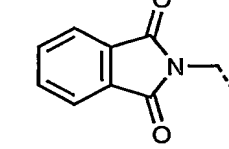
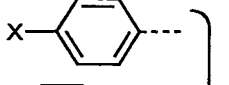
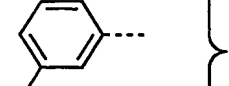
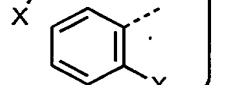
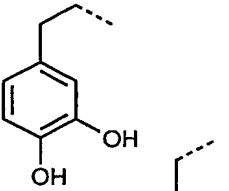
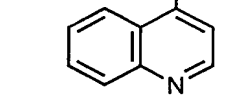
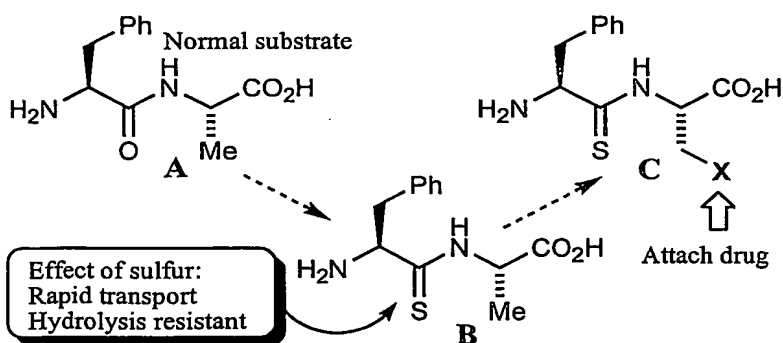
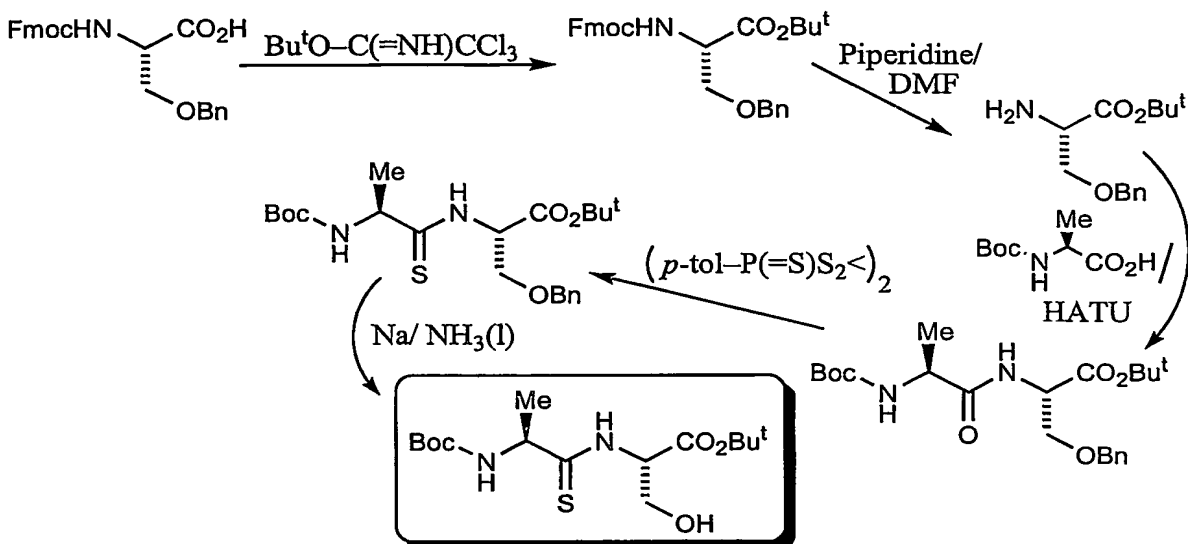


**Figure 1**

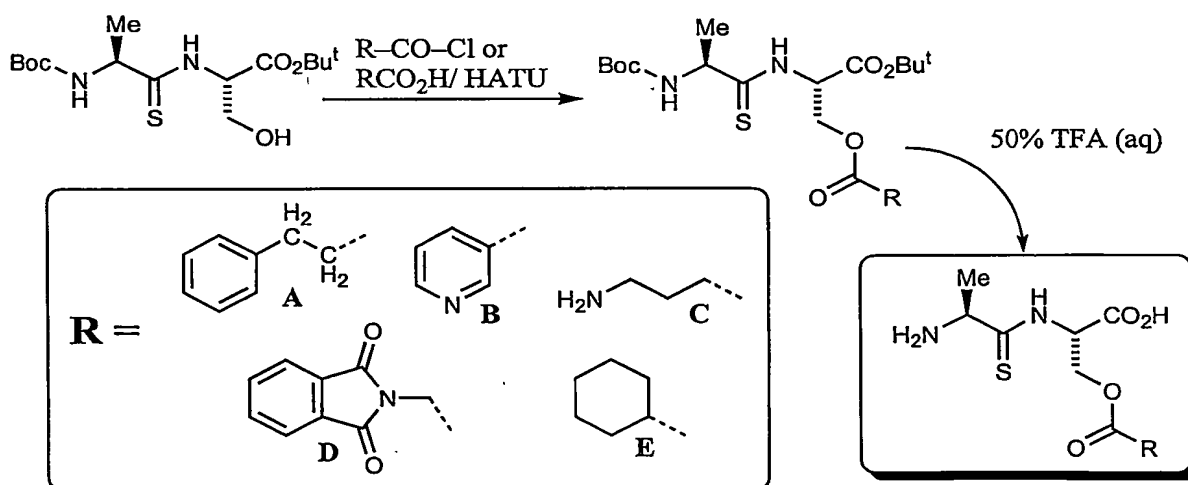
| Column A<br>Thiodipeptide core   | Column B<br>Y gps (cf. drugs)   | Column C<br>Y gps (cf. drugs)  | Column D<br>Side-chain R <sup>1</sup> gps   |
|--|---|--|---|
| <br> | <br><br><br><br> | <br><br><br><br> | H<br>Me<br>CH <sub>2</sub> Ph<br>CHMe <sub>2</sub><br>CH <sub>2</sub> OH<br>CH <sub>2</sub> SH<br>CH <sub>2</sub> CO <sub>2</sub> H<br>CH <sub>2</sub> CONH <sub>2</sub><br>(CH <sub>2</sub> ) <sub>4</sub> NH <sub>2</sub> |

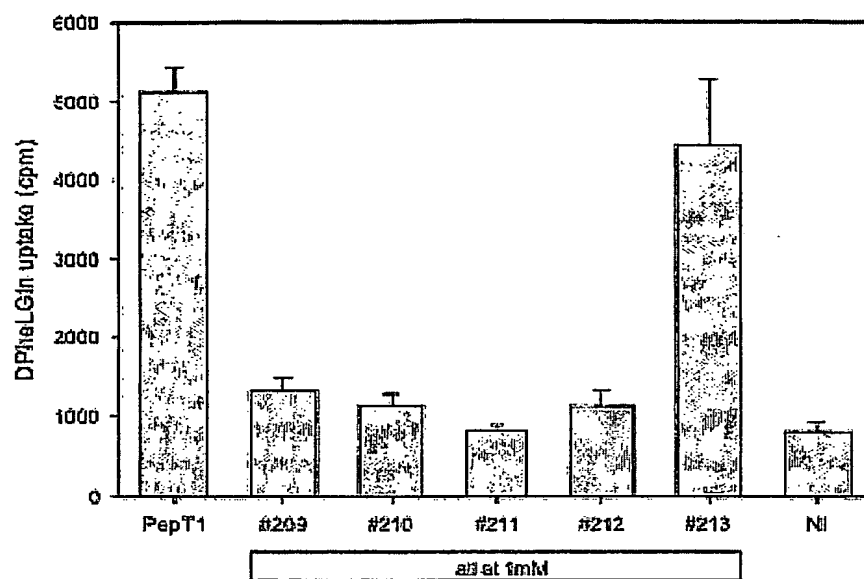
**Figure 2.**

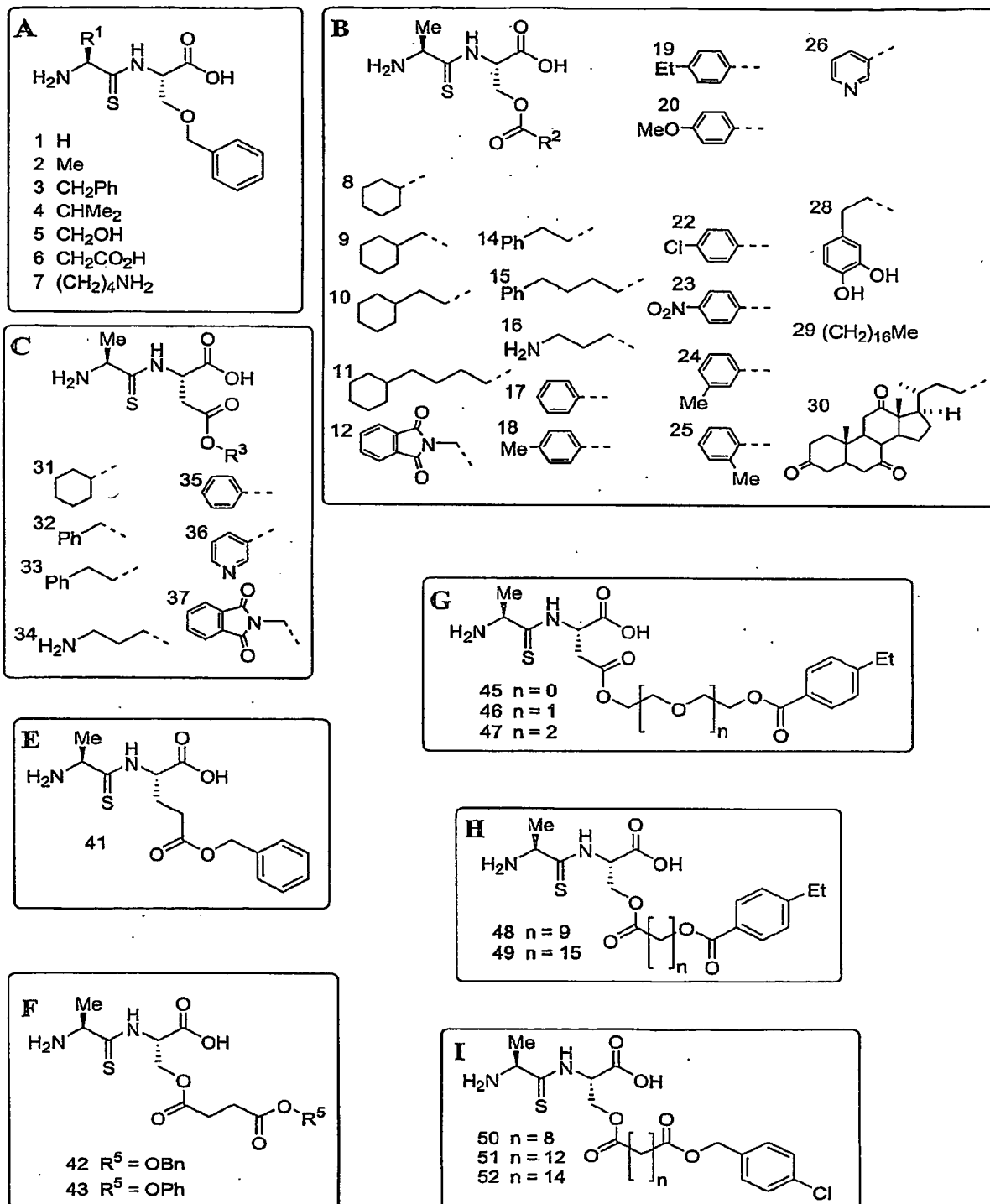
**Figure 3**



**Figure 4**



**Figure 5**

**Figure 6****Selected synthetic thiopeptide targets prepared as drug carrier models**

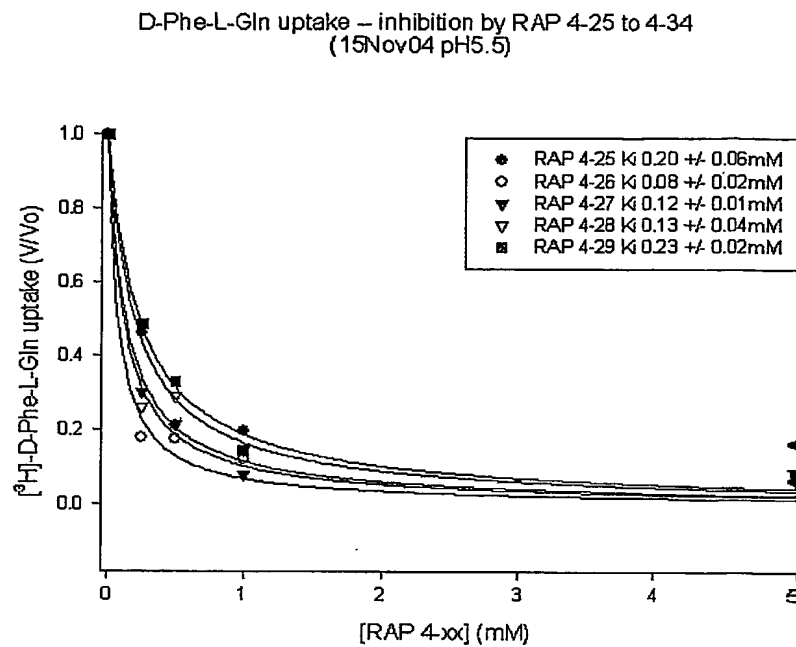
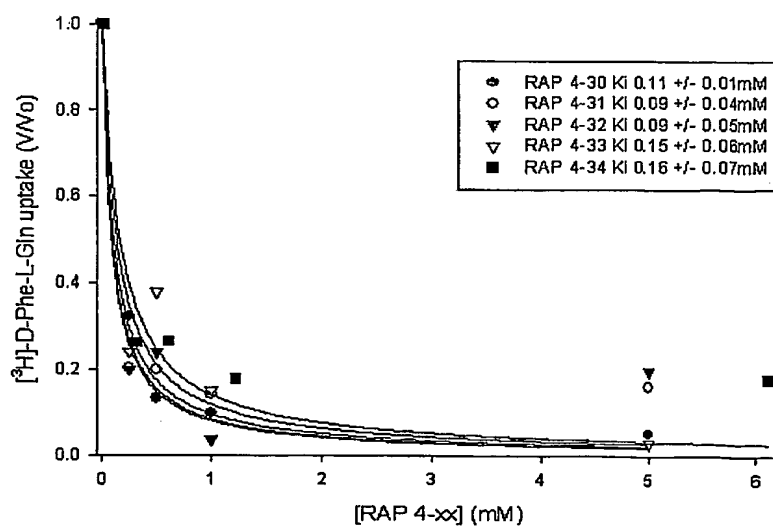
**Figure 7**

| Entry | Type | Binding<br>K <sub>i</sub> /mM | Transport<br>(cf Gly-Gln)† |
|-------|------|-------------------------------|----------------------------|
| 1     | A    | 1.41                          |                            |
| 2     | A    | 0.29                          |                            |
| 3     | A    | 0.37                          |                            |
| 4     | A    | 0.09                          |                            |
| 5     | A    | 0.34                          | <                          |
| 6     | A    | 0.53                          |                            |
| 7     | A    | 1.05                          |                            |
| 8     | B    | 0.3                           | =                          |
| 9     | B    | 0.2                           | =                          |
| 10    | B    | 0.12                          | =                          |
| 11    | B    | 0.11                          | =                          |
| 12    | B    | 0.03                          | <                          |
| 14    | B    | 0.1                           | <                          |
| 15    | B    | 0.09                          | =                          |
| 16    | B    | 0.6                           | <                          |
| 17    | B    | 0.23                          | <                          |
| 18    | B    | 0.2                           | =                          |
| 19    | B    | 0.13                          | <                          |
| 20    | B    | 0.16                          | =                          |
| 22    | B    | 0.09                          | >                          |
| 23    | B    | 0.11                          | =                          |
| 24    | B    | 0.08                          |                            |
| 25    | B    | 0.12                          | =                          |
| 26    | B    | 0.36                          | <                          |
| 28    | B    | 0.05                          | =                          |
| 29    | B    | n.d.*                         | <                          |
| 30    | B    | n.d.*                         | =                          |
| 31    | C    | 0.61                          |                            |
| 32    | C    | 0.25                          |                            |
| 33    | C    | 0.09                          |                            |
| 34    | C    | 1.58                          |                            |
| 35    | C    | 0.19                          | =                          |
| 36    | C    | 2.32                          | <                          |
| 37    | C    | 0.93                          |                            |
| 41    | E    | 0.03                          |                            |
| 42    | F    | 0.09                          | <                          |
| 43    | F    | 0.15                          | =                          |
| 45    | G    | 0.24                          | >                          |
| 46    | G    | 0.07                          | >                          |
| 47    | G    | 0.22                          | =                          |
| 48    | H    | 1.3                           | =                          |
| 49    | H    | 4.69                          | =                          |
| 50    | I    | 0.15                          | >                          |
| 51    | I    | 2.5                           |                            |
| 52    | I    | 9.7                           | =                          |

\* 29B/30B are strong inhibitors (est. K<sub>i</sub> < 1mM), but are too insoluble for accurate K<sub>i</sub> determination.

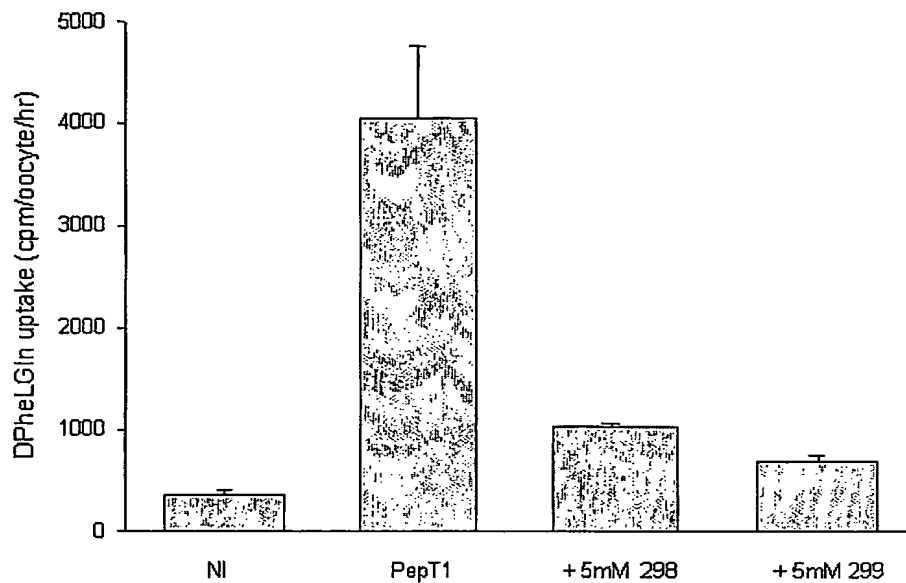
† Efflux of radio-labelled D-Phe-L-Gln is a positive indication of active transport via PepT1 (although negative results do not necessarily mean that substrates are not transported). In these assays (see elsewhere for details), the dipeptide Gly-Gln (which is known to be transported) caused *ca* 30% of labelled D-Phe-L-Gln to remain in the oocytes; at the same concentration, substrates were assessed as:

- < (measurable efflux, but less transport than Gly-Gln, with  $\geq 50\%$  of D-Phe-L-Gln remaining)
- = (efflux similar to the effect of Gly-Gln – i.e. 25-50% of labelled D-Phe-L-Gln remaining)
- > (more efflux than Gly-Gln – i.e. <25% of labelled D-Phe-L-Gln remaining)

**Figure 8a****Figure 8b**

**Figure 9a**

Inhibition of DPheLGln uptake by Rachel 298 &amp; 299 5mM15/7/04

**Figure 9b**

Efflux from PepT1-expressing oocytes, pH 5.5 90 min 2/8/04

